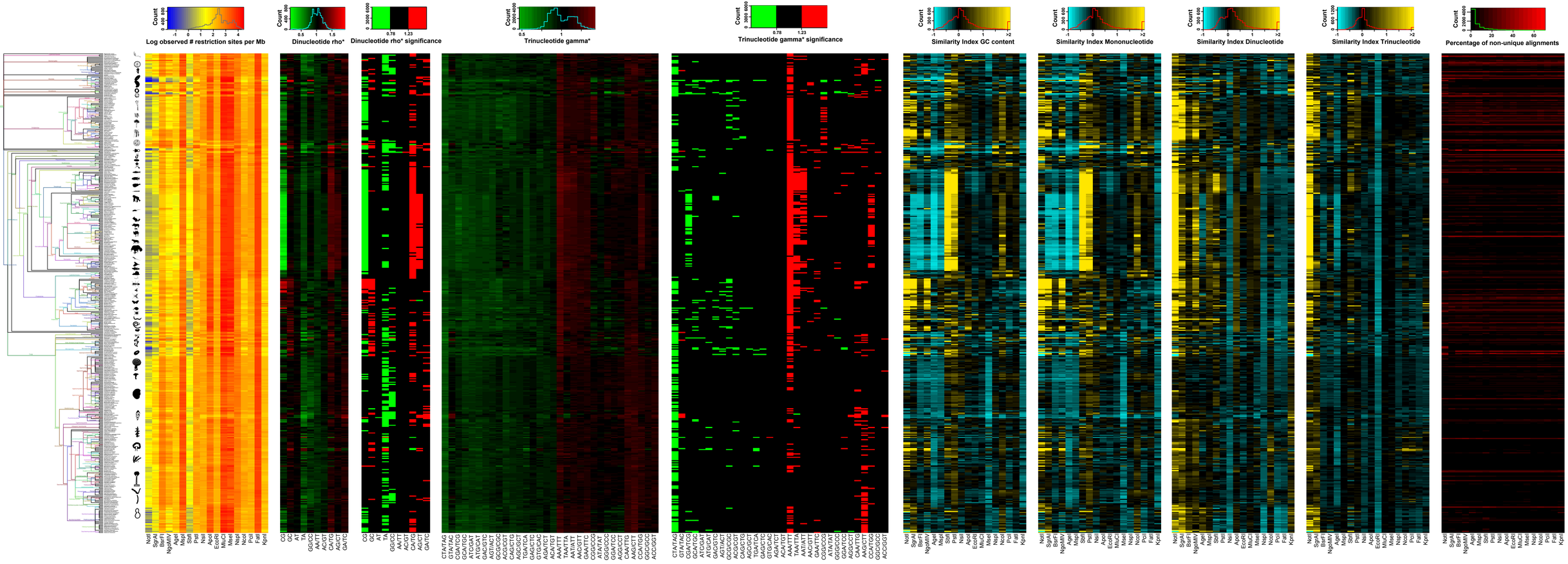
Log observed # restriction sites per Mb
Dinucleotide rho*
Dinucleotide rho* significance
Trinucleotide gamma*
Trinucleotide gamma* significance
Similarity Index GC content
Similarity Index Mononucleotide
Similarity Index Dinucleotide
Similarity Index Trinucleotide
Percentage of non-unique alignments
NotI SgrAI BsrFI NgoMIV AgeI MspI SbfI PstI NsiI ApoI EcoRI MluCI MseI NspI NcoI FspI FatI KpnI
CG GC AT TA GG/CC AA/TT AC/GT AT/AT AG/CT GA/TC
CTA/TAG GTA/TAC CGA/TCG GCA/TGC ATC/GAT ATG/CAT GAC/GTC AGT/ACT GCG/CGC AGC/GCT AGG/CCT TGA/TCA GAG/CTC GGC/GCC AGA/TCT ACA/TGT AAA/TTT TAA/TTA AAC/GTT GAA/TTC CGC/GCG ATA/TAT GGG/CCC AGA/TCC AGG/CCT CAA/TTG AAG/CTT CCA/TGG GCC/GGC ACC/GGT
ATA/TAT CGC/GCG AAG/CTT GGA/TCC AGG/CCT CAA/TTG CCA/TGG GCC/GGC GGG/CCC ACC/GGT